

## **CHEMOKINE PARC SUPPRESSES SPECIFIC CYTOKINE PRODUCTION**

### **Field of the Invention**

- [01] The invention relates to the field of proliferative cell disorders. In particular, it relates to the cytokines stimulated by chemokine PARC in leukocytes.

### **Background of the Prior Art**

- [02] Ordered arrays of proteins provide an attractive strategy for high-throughput analysis of proteins. To be truly useful for this purpose, however, such arrays must yield sensitive, quantitative, and reproducible measurements of protein levels. It is also desirable that assays on these arrays utilize small sample volumes and be amenable to automated systems for high-throughput processing. There have been a number of recent examples of the use of protein arrays for a variety of applications (1-6). While these approaches have established the feasibility of protein arrays, they have not yet demonstrated practical utility for measuring protein expression levels in a manner analogous to a gene expression array. A microarray consisting of immobilized antibodies is the most straightforward near-term approach for developing a chip for highly parallel analysis of protein levels. Experience with such arrays is limited, and the levels of sensitivity (ca. 10 ng/mL) and multiplexing have been insufficient for quantifying most biological change (7-10).
- [03] PARC (pulmonary and activation-regulated chemokine) is also known as AMAC-1 (alternative activated macrophage associated C-C-chemokine ), MIP-4 (macrophage inflammatory protein-4 ), DC-CK1 (dendritic cell-derived chemokine-1) and CCL18. It is expressed mainly by dendritic cells (DC) and macrophages, and is chemotactic mainly for T lymphocytes. PARC is expressed in alveolar macrophages (J Immunol 1997 Aug 1;159(3):1140-9), follicular dendritic cells and macrophages (J Leukoc Biol 2001 May;69(5):785-93). PARC has been shown to be chemotactic in vivo for both

CD4(+) and CD8(+) T lymphocytes and may therefore be implicated in both humoral and cell-mediated immunity (Genomics 1999 Mar 15;56(3):296-302).

- [04] PARC is mainly induced by Th2-associated cytokines and inhibited by Th1-associated cytokines. PARC is specifically induced in macrophages by cytokines such as, IL-4, IL-13, and IL-10 (mediators of Th2 response). Expression of PARC is inhibited by IFN-gamma (mediator of Th1 response). Peripheral blood monocytes do not express PARC; time course experiments show that monocyte-to-macrophage differentiation is a prerequisite for PARC expression. In vivo, PARC is expressed by alveolar macrophages from healthy persons, smokers, and asthmatic patients. PARC expression is induced by Th2-associated cytokines; thus, PARC may be involved in the APC-dependent T cell development in inflammatory and immune reactions (J Immunol 1998 Feb 1;160(3):1411-8). The Th1/Th2 paradigm underlying T-cell polarization has been extended to the corresponding concept of APC polarization. Macrophages as well as DCs can undergo Th1- or Th2-like polarization; APC1 and APC2 thus acquire the capacity to drive the development of naive T cells and the reactivation of resting T cells towards either a Th1 or a Th2 phenotype, respectively. APC1 are classically activated by mediators such as IFN-gamma, TNF-alpha or LPS, while APC2 are alternatively activated by IL-4, IL-10 or PGE. (Skin Pharmacol Appl Skin Physiol 2001 Sep-Oct;14(5):272-9. PARC expression is induced by Th2-associated cytokines such as IL-4, IL-13 and IL-10, and is inhibited by Th1-associated cytokines such as IFN-gamma. The Th2-associated expression pattern of PARC in alternatively activated suppressor macrophages in vivo and in vitro and its absence from epidermal Langerhans cells in vivo suggest that it may be involved in inhibition of Th1 reactions and in tolerance induction (Res Immunol 1998 Sep-Oct;149(7-8):633-7).

- [05] Interestingly, there is a suggestion in the art of a potential application of PARC and its "derivatives" in the treatment of allergy. Allergic reactions are characterized by the infiltration of tissues by activated eosinophils, Th2 lymphocytes, and basophils. The

beta-chemokine receptor CCR3, which recognizes the ligands eotaxin, eotaxin-2, MCP-3, MCP-4, and RANTES, plays a central role in this process. PARC, which is known to act as a T cell chemoattractant, exhibits CCR3 antagonistic activity. A modified form of PARC, called Met-chemokine beta 7 (Met-Ckbeta7), is a potent and specific CCR3 antagonist that prevents signaling through this receptor and, at concentrations as low as 1 nM, can block eosinophil chemotaxis induced by the most potent CCR3 ligands. The enhanced activity of Met-Ckbeta7 is due to the alteration of the extreme N-terminal residue from an alanine to a methionine. Met-Ckbeta7 is a more potent CCR3 antagonist than Met- and aminooxypentane (AOP)-RANTES and, unlike these proteins, exhibits no partial agonist activity and is highly specific for CCR3. PARC may therefore use chemokine receptor agonism and antagonism to control leukocyte movement in vivo. Thus, this antagonist and its "derivatives" may be of use in ameliorating leukocyte infiltration associated with allergic inflammation and they have potential therapeutic use in the treatment of allergy (J Immunol 2000 Feb 1;164(3):1488-97).

- [06] In a study of macrophage- and T-lymphocyte-specific chemoattractants involved in the positioning of immune effector cells during the elicitation phase of contact hypersensitivity (a prototype of a T-lymphocyte-mediated immune reaction), it has been demonstrated that expression of the T-lymphocyte-specific chemokines IP-10 and MIG in epidermis and dermis, and of MDC, PARC and TARC exclusively in the dermis started after 12 hours reaching maximum levels at 72 hours and was associated with infiltration of T cells into the epidermal compartment (Am J Pathol 2001 Feb;158(2):431-40). PARC expression was significantly increased in lungs affected by hypersensitivity pneumonitis (HP), a lung inflammatory disorder characterized by accumulation of T lymphocytes, in comparison with lungs affected by idiopathic pulmonary fibrosis and controls. Macrophages, DCs, and alveolar epithelial cells were the main sources of PARC. There was a direct correlation between the levels of tissue PARC and the number of lymphocytes in the bronchoalveolar lavage fluids. High levels of PARC were detected in the subacute rather than the chronic phase of HP.

These findings suggest a role for PARC in the pathogenesis of HP (J Leukoc Biol 2001 Oct;70(4):610-6).

- [07] Gene expression of PARC and ELC but not of LARC or SLC has been found in human atherosclerotic plaques, where PARC mRNA was restricted to CD68+ macrophages. The expression patterns of PARC and ELC mRNA in human atherosclerotic lesions suggest a potential role for these two CC chemokines in attracting T lymphocytes into atherosclerotic lesions. (Am J Pathol 1999 Feb; 154(2):365-74).
  
- [08] In livers of patients with chronic hepatitis C, PARC expression has been detected in mononuclear cells mainly in the portal area. The mRNA levels significantly correlated with serum alanine aminotransferase levels ( $p < 0.001$ ). These results suggest that the local production of PARC and RANTES participates in immune responses by attracting naive and active T cells to the portal and periportal areas, respectively (Lab Invest 2000 Mar;80(3):415-22).
  
- [09] CCL18/PARC is selectively induced by staphylococcal enterotoxins in mononuclear cells and PARC levels are enhanced in septic and rheumatoid arthritis. CCL18/PARC protein is spontaneously secreted by PBMC and is selectively induced in PBMC by staphylococcal enterotoxins (SEA, SEB) and IL-4, but not by IFN-gamma and the CXCL8/IL-8 inducers LPS or Concanavalin A. In synovial fluids from septic and rheumatoid arthritis patients, four fold-enhanced CCL18/PARC levels (150 ng/ml) were detected compared to those in crystal-induced arthritis and osteoarthritis. In septic arthritis, the synovial levels of CCL18/PARC were fivefold higher than those of CXCL8/IL-8. Immunohistochemistry revealed CD68(+) monocytes/macrophages as the main CCL18/PARC-producing cell type in both PBMC and arthritic synovial tissue. In addition, CD1a(+) blood dendritic cells expressed CCL18/PARC. These findings

suggest that monocytic cells respond to Gram-positive bacterial infection by the production of CCL18/PARC in the synovial cavity (Eur J Immunol 2001 Dec;31(12):3755-62 PMID: 11745396).

- [10] Using cDNA microarrays, elevated expression of chemokines MIP-1beta and PARC was observed in preterm labour with chorioamnionitis (PTL(+INF)) compared to PTL(-INF) amnion and choriondecidua respectively. Likewise, the cytokines oncostatin-M and PBEF were more highly expressed in PTL(+INF) compared with PTL(-INF) and in spontaneous labour at term compared with Caesarean section at term, respectively. Conversely, inhibin A, TIMP-3 and TIMP-4 were all significantly elevated in PTL(-INF) compared with PTL(+INF). Furthermore, differential expression patterns of classes of genes, grouped according to function (e.g. chemokines), were noted (Mol Hum Reprod 2002 Apr;8(4):399-408 PMID: 11912289).
  
- [11] In ascitic fluids from patients with ovarian carcinoma, significantly higher levels of CXCL8 and CCL18 (PARC) were detected compared with those in nonovarian carcinoma patients. In contrast to CXCL8, CCL18 was not inducible in carcinoma cell lines. Immunostaining showed CCL18 expression in tumor-infiltrating cells with monocyte/macrophage morphology but not in the ovarian carcinoma cells. Biochemically heterogeneous but biologically active forms of several chemokines are present at different concentrations in ovarian carcinoma ascitic fluid. This points to a delicate balance of chemokines in epithelial ovarian cancer and to a potentially major role for CXCL8 and CCL18 in this tumor (J Biol Chem 2002 Jul 5;277(27):24584-93 PMID: 11978786).
  
- [12] Cluster analysis of DNA microarray profile of differentially expressed genes in a monkey model of allergic asthma revealed at least five groups of genes with unique

expression patterns. One cluster contained genes for several chemokine mediators including eotaxin, PARC, MCP-1 and MCP-3 (Genome Biol 2002;3(5):research0020 PMID: 12049661).

- [13] There is a need in the art for improved diagnostic and therapeutic techniques for diseases that are associated with inflammatory processes.

#### **BRIEF SUMMARY OF THE INVENTION**

- [14] According to one embodiment of the invention a method is provided for treating a disease characterized by an undesirable amount of cell proliferation. An effective amount of chemokine PARC is administered to a patient who has a proliferative cell disorder, whereby undesirable proliferation of cells is reduced.
- [15] According to another embodiment of the invention a method of diagnosing a proliferative cell disorder in a patient is provided. The amount of PARC is determined in a test sample from a patient. The amount determined is compared to an average amount found in a control sample of a population of healthy humans. A decreased amount in the test sample relative to the average amount indicates a proliferative cell disorder in the patient.
- [16] According to still another embodiment of the invention a method of stimulating mitosis is provided. An inhibitory molecule for chemokine PARC is administered to a subject in need of augmented mitosis. The inhibitory molecule is selected from the group consisting of antibodies specific for chemokine PARC and antisense molecules which bind to and inhibit transcription of PARC mRNA. Mitosis in the subject is thereby increased.
- [17] A still further embodiment of the invention provides a method to monitor the effects of PARC therapy or anti-PARC therapy. The amount of one or more proteins in a

sample collected from a patient at a first time is determined. The protein is selected from the group consisting of AR, IL-1sR1, IL-2, IL-6sR, LIF, SDF-1a, and fragments thereof. The determination is repeated on a sample collected from the patient at a later time. The amounts measured at the two times is compared. A decreased amount over time denotes an effect of PARC and a increased amount denotes an effect of an anti-PARC inhibitory molecule.

- [18] The invention thus provides the art with useful diagnostic and therapeutic methods for diseases characterized by excess mitosis.

#### **BRIEF DESCRIPTION OF THE DRAWINGS**

- [19] Figure 1. *Cartoon of immunoassays with RCA signal amplification:*
- [20] (A). In the adaptation of RCA used for protein signal amplification, the 5' end of an oligonucleotide primer is attached to an antibody. (B) The antibody-DNA conjugate binds to its specific target molecule; in the multiplexed microarray immunoassay, the targets are biotinylated secondary antibodies and the conjugate is an antibiotin antibody. (C) A circular DNA molecule hybridizes to its complementary primer on the conjugate, and in the presence of DNA polymerase, and nucleotides, rolling circle replication occurs. (D) A long single DNA molecule that represents a concatamer of complements of the circle DNA sequence is generated that remains attached to the antibody. (E) This RCA product is detected by hybridization of multiple fluorescent, complementary oligonucleotide probes. RCA product fluorescence is measured with a conventional microarray scanning device. The amount of fluorescence at each spot is directly proportional to the amount of specific protein in the original sample.
- [21] Figure 2. *Cytokine/Chemokines affected by treatment of CD4+ cells.*
- [22] CD4+ cells were treated for 48 hours with the substance indicated on the x-axis. PARC had a suppressive effect.

- [23] Figure 3. *Selected Cytokine/Chemokines affected by PARC.*
- [24] Conditions were as described in the brief description of Figure 2.

#### **DETAILED DESCRIPTION OF THE INVENTION**

- [25] It is a discovery of the present inventors that chemokine PARC suppresses expression of certain sets of cytokines and chemokines in human leukocytes. Based on the function of these induced cytokines, it is suggested that PARC has novel anti-mitotic activity.
- [26] Diseases and conditions that can be advantageously diagnosed and treated according to the present invention include without limitation autoimmune disease, rheumatoid arthritis, cancer, myeloproliferative disease, and coronary artery disease. Other such syndromes are also amenable to the methods of the invention.
- [27] Test samples used for performing the diagnostic method are preferably from serum, plasma, blood, lymph fluid, peripheral lymphatic tissue, or blood. Desirably the test sample contains, or has contained, leukocytes, monocytes, dendritic cells, or Langerhans cells. However, it may be desirable that the actual sample upon which the assay is performed be relatively free of cells.
- [28] Altered expression of a cytokine can be determined relative to a control sample. The control sample can be obtained from an organ distal to the area of excess mitosis in the test subject. Alternatively, the control sample can be obtained from a subject or subjects not experiencing or evidencing any such disorder. An average value or range can be determined from a population of healthy individuals and used as a control value. Altered expression can be determined at any threshold that is statistically significant. This can be an decrease relative to a control sample of 25%, 50%, or 75%, for example. The threshold can be set to at least one tenth the level of the control



sample. Alternatively, the threshold can be set to at least one one-hundredth the level in the control sample.

[29] Altered expression of a cytokine can be determined using either mRNA or protein as an indication of expression level. Preferably the protein will be determined. The determination need not be strictly quantitative. For example, in cases where a cytokine goes from an unexpressed to an expressed state a qualitative assessment may be sufficient. Any assay known in the art for detecting gene expression can be used, either individually or multiplexed. The assays used may involve gene arrays, protein arrays, antibody arrays, Western blotting, ELISA, immunoprecipitation, filter binding assays, hybridization assays, etc. The protein microarray employing a rolling circle amplification for detection described in detail below is preferred, but need not be used. Briefly, capture antibodies are affixed to a solid support in a predetermined pattern (array) and test sample is applied to the array so that proteins (cytokines) in the test sample can bind to antibodies on the array which are specific for that particular protein. Second antibodies are applied which are specific for the same set of proteins as are the capture antibodies. The second set of antibodies can be labeled with a hapten. A third set of antibodies is then applied to the array. The third set of antibodies is specific for the hapten on the second set of antibodies or with the constant region of the second set of antibodies. The third set of antibodies contains an attached oligonucleotide. The oligonucleotide can be used as a primer to amplify a template to create an amplification signal. Preferably the template is a circular DNA such that rolling circle amplification can create a large signal. Alternatively, the second antibody can be directly detectable, for example by rolling circle amplification of an attached oligonucleotide.

[30] If increased mitosis is desired, then a subject can be treated by administering an antibody which specifically binds to a human PARC. The antibody can be a monoclonal or polyclonal antibody. It can be a complete antibody molecule or a fragment. Standard antibody fragments are known in the art and any of these can be used, including Fab, F(ab')<sub>2</sub>. Single chain Fv (ScFv) can also be used. The antibodies can if desired be attached to other moieties, such as therapeutic agents. Single

antibodies or cocktails of antibodies can be used. The cocktails can be directed to the same or different cytokines. Antibodies can be administered by any means known in the art, including but not limited to intravenous, intrathecal, directly to the thymus or to a lymph nodes, subcutaneous, oral, and intramuscular. Antisense molecules can also be used which specifically bind to mRNA encoding human PARC and inhibit expression of an PARC.

- [31] PARC-treated leukocytes (CD34<sup>+</sup> cell and monocytes) expressed decreased amounts of a set of cytokine, chemokines, growth factors, and soluble receptors, including AR (amphiregulin), IL-1sR1 (IL-2 soluble receptor 1), IL-2, IL-6sR (IL-6 soluble receptor), LIF(Leukocyte inhibitory factor), SDF-1a (Stromal cell-derived factor-1-alpha), BDNF (brain-derived neurotrophic factor, FGF-6(fibroblast growth factor-6), Flt-3 ligand (fms-like tyrosine kinase-3 ligand), G-CSF(Granulocyte colony stimulating factor), HCC4 (hemofiltrate CC chemokine 4), IL-12p70.
  
- [32] PARC is a therapeutic target. Inhibitors (in the form of antibodies, small molecular drugs, anti-sense RNA therapy) of PARC can be used to treat diseases associated with too little mitosis. PARC and antibodies and anti-sense molecules thereto can be administered by any technique known in the art. Such methods include, but are not limited to intravenous, intramuscular, subcutaneous, oral, nasal and intrabronchial injections or instillations. Therapeutics based on inhibition of PARC can take the form of proteins, antibody-based therapy or small molecular drugs, anti-sense RNA therapies. The receptors for PARC can also be considered as a therapeutic target for protein therapy, antibody therapy or small molecular drug therapy.
  
- [33] PARC inhibitory molecules can be administered to other individuals in need of an increased mitosis. Such a situation might occur, for example, in the case of wound healing. We utilized highly sensitive antibody based microarray protein chips, which detect 78 cytokines/chemokines simultaneously. Our result demonstrated that a specific set of cytokines/chemokines was suppressed in PARC-treated immature dendritic cells, monocytes and a monocyte cell line.

## EXAMPLES

### Example 1--Materials and Methods

[34] In Experiment 1, 16 cell culture supernatants (RPMI supplemented with GMCSF and IL-4) were provided by Drs. De Yung and Zack Howard of NCI (NCI Frederick, Frederick, MD 21702). The cells had been treated as follows:

1. Medium alone without cells, a background control.
2. Medium with cells, a negative control.
3. LARC at 100 ng/ml, human chemokine.
4. hBD2 at 1000 ng/ml, human beta defensin 2.
5. hBD3 at 1000 ng/ml, human beta defensin 3.
6. PARC at 1000 ng/ml, human chemokine.
7. hNPm at 1000 ng/ml, natural human neutrophil defensins (a), mixture of hNP1, hNP2 and hNP3 and isolated from the granules of polymorphonuclear leukocyte.
8. hNP1 at 1000 ng/ml, human neutrophil protein, alpha defensin.
9. hEDN at 1000 ng/ml, human eosinophil derived neurotoxin.
10. mEAR2 at 1000 ng/ml, mouse protein, no effect on human cells and is a negative control.
11. RNase1 at 1000 ng/ml, human RNase 1, eosinophil derived, It can strongly activate iDC and is a control for iDC maturation.
12. C5a at 10 nM, complement factor 5a.
13. W pep. at 100 nM, hexapeptide.
14. PAF at 10 ng/ml, platelet activating factor.
15. RANTES at 100 ng/ml, human chemokine.
16. TNFa at 50 ng/ml, a positive control.

[35] In Experiment 2, 84 cell culture supernatants (RPMI supplemented with GMCSF and IL-4) were provided by Dr. De Yung and Dr. Zack Howard of NCI (NCI Frederick, Frederick, MD 21702). Samples were divided into following 5 groups:

1. Group 1: (Time-course, 36 samples) monocyte-derived DCs and CD34-derived DCs treated with RNase 1, hEDN (Rnase 2) or RNase 3 for the following times: 0, 2 or 3, 6, 12, 24, and 48 hours.
2. Group 2: (Concentration-dependence, 29 samples) monocyte-derived DCs and CD34-derived DCs treated for 48 hrs with 10, 100, 500, or 1000 ng/ml of RNase 1 or hEDN (Rnase 2); or with 1000 or 3000 ng/ml of RNase3.
3. Group 3: (RNase activity-dependence, 6 samples) CD34-derived DCs treated with 1000 ng/ml RNase 1 or 2 in the presence of ribonuclease inhibitor.
4. Group 4: (Cell type specificity, 8 samples) lymphocytes treated with RNase 1 or hEDN (Rnase 2). Monocyte cell lines treated with RNase 1, hEDN (Rnase 2) or RNase 3.
5. Group 5: (independent RNase source, 5 samples) Monocytes treated with 1000 ng/ml RNase 1, hEDN (Rnase 2) or RNase 3.

[36] **Microarray manufacture:** Antibody microarrays were printed using a Packard Biosciences (Downers Grove, IL) BCA-II piezoelectric microarray dispenser on cyanosilane-coated glass slides divided by Teflon boundaries into sixteen 0.5 cm diameter circular subarrays. Monoclonal antibodies (commercially available) for 78 cytokines were dispensed in quadruplicate at a concentration of 0.5 mg/ml. Printed slides were blocked as described [21] and stored at 4°C until use. Batches of slides were subjected to a quality control consisting of incubation with a fluorescently-labeled anti-mouse antibody, followed by washing, scanning and quantitation. Typically, the coefficient of variability (CV) of antibody deposition in printing was <5%.

[37] **RCA Immunoassay:** The assay was performed by a liquid-handling robot (Biomek 2000, Beckman Instruments, Fullerton, CA, which was enclosed in an 80% humidified, HEPA-filtered, plexiglass chamber. For each sample, duplicates were

tested either neat or diluted 1:10. 20 µl of samples was applied to each sub-array and immunoassays with RCA signal amplification were performed as described [21] Slides were scanned (GenePix, Axon Instruments Inc., Foster City, CA) at 10-µm resolution with laser setting of 100 and PMT setting of 550. Mean pixel fluorescence were quantified using the fixed circle method in GenePix Pro 3.0 (Axon Instruments, Foster City, CA). The fluorescence intensity of 8 microarray features (duplicates subarrays and quadruplicates spots in each subarray) was averaged for each feature and sample, and the resulting cytokine values were determined. For every slide, a set of blanks was run as a negative control.

- [38] ***Data Quality Control:*** Subarray(s) were excluded from analysis if fluorescent intensities were generally weak (indicating weak RCA in that particular subarray), if there were visible defects in the array (such as scratches), or if there was high background signal. A total of 168 subarrays (84 samples) were analyzed, 4 subarray-1s and 4 subarray-2s were excluded on the basis of these quality control criteria. None of the samples failed both duplicated subarrays. Analyses were performed using complete set of data containing the levels of all 78 cytokines from 84 cell culture supernatants. Untransformed fluorescent intensities were used as data values in all of the analyses.

#### Example 2—Suppression by PARC

- [39] The results of Experiment 1, described in Example 1 above, measured at 48 hours are shown in Figures 2 and 3. These data demonstrate that the chemokine PARC suppresses expression of a set of cytokines including AR, IL-1sR1, IL-2, IL-6sR, LIF, AND SDF1a. Therefore PARC has utility in diseases characterized by too much mitosis, such as coronary artery disease, rheumatoid arthritis, cancers, autoimmune diseases and myeloproliferative disorders.

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